

Input Set : A:\SeqListing.txt

```
3 <110> APPLICANT: ImClone Systems Incorporated
W--> 4 <120> TITLE OF INVENTION: Human Antibodies Specific To KDR And Uses Thereof
W--> 5 <130> FILE REFERENCE: 11245/47802
W--> 6 <140> CURRENT APPLICATION NUMBER: To Be Assigned
C--> 7 <141> CURRENT FILING DATE: 2004-09-04
     8 <150> PRIOR APPLICATION NUMBER: PCT/US03/06459
     9 <151> PRIOR FILING DATE: 2003-03-04
    10 <150> PRIOR APPLICATION NUMBER: 60/361,783
    11 <151> PRIOR FILING DATE: 2002-03-04
W--> 12 <160> NUMBER OF SEQ ID: 93
    13 <170> SOFTWARE: WordPerfect 8.0 for Windows
    16 <210> SEO ID NO: 1
    17 <211> LENGTH: 11
    18 <212> TYPE: PRT
    19 <213> ORGANISM: Human
    21 <400> SEQUENCE: 1
    23
         Arg Ala Ser Gln Ser Val Ser Ser Tyr Leu Ala
    24
                         5
    26 <210> SEQ ID NO: 2
    27 <211> LENGTH: 7
    28 <212> TYPE: PRT
    29 <213> ORGANISM: Human
    31 <400> SEQUENCE: 2
        Asp Ser Ser Asn Arg Ala Thr
    34
    36 <210> SEQ ID NO: 3
    37 <211> LENGTH: 9
    38 <212> TYPE: PRT
    39 <213> ORGANISM: Human
    41 <400> SEQUENCE: 3
        Leu Gln His Asn Thr Phe Pro Pro Thr
    46 <210> SEQ ID NO: 4
    47 <211> LENGTH: 11
    48 <212> TYPE: PRT
    49 <213> ORGANISM: Human
    51 <400> SEQUENCE: 4
    53
        Arg Ala Ser Gln Gly Ile Ser Ser Arg Leu Ala
    56 <210> SEQ ID NO: 5
    57 <211> LENGTH: 7
    58 <212> TYPE: PRT
    59 <213> ORGANISM: Human
```

Input Set : A:\SeqListing.txt
Output Set: N:\CRF4\09152004\J506997.raw

```
61 <400> SEOUENCE: 5
   Ala Ala Ser Ser Leu Gln Thr
63
64
66 <210> SEQ ID NO: 6
67 <211> LENGTH: 9
68 <212> TYPE: PRT
69 <213> ORGANISM: Human
71 <400> SEQUENCE: 6
   Gln Gln Ala Asn Arg Phe Pro Pro Thr
76 <210> SEQ ID NO: 7
77 <211> LENGTH: 14
78 <212> TYPE: PRT
79 <213> ORGANISM: Human
81 <400> SEQUENCE: 7
     Ala Gly Thr Thr Thr Asp Leu Thr Tyr Tyr Asp Leu Val Ser
84
86 <210> SEQ ID NO: 8
87 <211> LENGTH: 7
88 <212> TYPE: PRT
89 <213> ORGANISM: Human
91 <400> SEQUENCE: 8
   Asp Gly Asn Lys Arg Pro Ser
96 <210> SEQ ID NO: 9
97 <211> LENGTH: 10
98 <212> TYPE: PRT
99 <213> ORGANISM: Human
101 <400> SEQUENCE: 9
103
    Asn Ser Tyr Val Ser Ser Arg Phe Tyr Val
104
106 <210> SEQ ID NO: 10
107 <211> LENGTH: 13
108 <212> TYPE: PRT
109 <213> ORGANISM: Human
111 <400> SEQUENCE: 10
113
      Ser Gly Ser Thr Ser Asn Ile Gly Thr Asn Thr Ala Asn
114
116 <210> SEQ ID NO: 11
117 <211> LENGTH: 7
118 <212> TYPE: PRT
119 <213> ORGANISM: Human
121 <400> SEQUENCE: 11
     Asn Asn Asn Gln Arg Pro Ser
124
126 <210> SEQ ID NO: 12
127 <211> LENGTH: 12
128 <212> TYPE: PRT
129 <213> ORGANISM: Human
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```
131 <400> SEQUENCE: 12
133
     Ala Ala Trp Asp Asp Ser Leu Asn Gly His Trp Val
134
                        5
136 <210> SEQ ID NO: 13
137 <211> LENGTH: 10
138 <212> TYPE: PRT
139 <213> ORGANISM: Human
141 <400> SEQUENCE: 13
143
    Gly Phe Thr Phe Ser Ser Tyr Ser Met Asn
144
146 <210> SEQ ID NO: 14
147 <211> LENGTH: 17
148 <212> TYPE: PRT
149 <213> ORGANISM: Human
151 <400> SEQUENCE: 14
    Ser Ile Ser Ser Ser Ser Tyr Ile Tyr Tyr Ala Asp Ser Val Lys
154
                        5
                                           10
156
     Gly
157
     17
159 <210> SEQ ID NO: 15
160 <211> LENGTH: 7
161 <212> TYPE: PRT
162 <213> ORGANISM: Human
164 <400> SEQUENCE: 15
    Val Thr Asp Ala Phe Asp Ile
167
169 <210> SEQ ID NO: 16
170 <211> LENGTH: 10
171. <212> TYPE: PRT
172 <213> ORGANISM: Human
174 <400> SEQUENCE: 16
176
    Gly Gly Thr Phe Ser Ser Tyr Ala Ile Ser
177
179 <210> SEQ ID NO: 17
180 <211> LENGTH: 18
181 <212> TYPE: PRT
182 <213> ORGANISM: Human
184 <400> SEOUENCE: 17
186
     Gly Gly Ile Ile Pro Ile Phe Gly Thr Ala Asn Tyr Ala Gln Lys Phe
187
     Gln Gly
189
190
       18
192 <210> SEQ ID NO: 18
193 <211> LENGTH: 16
194 <212> TYPE: PRT
195 <213> ORGANISM: Human
197 <400> SEQUENCE: 18
199
     Gly Tyr Asp Tyr Tyr Asp Ser Ser Gly Val Ala Ser Pro Phe Asp Tyr
200
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Input Set : A:\SeqListing.txt

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202 <210> SEQ ID NO: 19
203 <211> LENGTH: 375
204 <212> TYPE: DNA
205 <213> ORGANISM: Human
207 <400> SEQUENCE: 19
209
      gag gtc cag ctg gtg cag tct ggg gct gag gtg aag aag cct ggg gcc
                                                                            48
210
      Glu Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
211
213
      tca gtg aag gtc tcc tgc aag gct tct gga ggc acc ttc agc agc tat
                                                                            96
214
      Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Thr Phe Ser Ser Tyr
215
                                        25
217
      get ate age tgg gtg ega eag gee eet gga eaa ggg ett gag tgg atg
                                                                           144
218
      Ala Ile Ser Trp Val Arq Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
219
                                   40
221
      gga ggg atc atc cct atc ttt ggt aca gca aac tac gca cag aag ttc
                                                                           192
222
      Gly Gly Ile Ile Pro Ile Phe Gly Thr Ala Asn Tyr Ala Gln Lys Phe
223
225
      cag ggc aga gtc act ttt acc gcg gac aaa tcc acg agt aca gcc tat
                                                                           240
226
      Gln Gly Arg Val Thr Phe Thr Ala Asp Lys Ser Thr Ser Thr Ala Tyr
227
                           70
                                                75
229
      atg gag ttg agg agc ctg aga tct gac gac acg gcc gtg tat tac tgt
                                                                           288
230
      Met Glu Leu Arg Ser Leu Arg Ser Asp Asp Thr Ala Val Tyr Tyr Cys
231
                       85
                                            90
233
      geg aga gga tac gat tac tat gat agt agt ggc gtg gct tcc ccc ttt
                                                                           336
      Ala Arg Gly Tyr Asp Tyr Tyr Asp Ser Ser Gly Val Ala Ser Pro Phe
234
235
                  100
                                      105
237
      gac tac tgg ggc cag gga acc ctg gtc acc gtc tca agc
                                                                           375
238
      Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
241 <210> SEQ ID NO: 20
242 <211> LENGTH: 125
243 <212> TYPE: PRT
244 <213> ORGANISM: Human
246 <400> SEQUENCE: 20
248
      Glu Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
249
251
      Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Thr Phe Ser Ser Tyr
252
                                        25
254
      Ala Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
255
                                    40
257
      Gly Gly Ile Ile Pro Ile Phe Gly Thr Ala Asn Tyr Ala Gln Lys Phe
258
                               55
                                                    60
260
      Gln Gly Arg Val Thr Phe Thr Ala Asp Lys Ser Thr Ser Thr Ala Tyr
261
263
      Met Glu Leu Arg Ser Leu Arg Ser Asp Asp Thr Ala Val Tyr Tyr Cys
264
                                            90
266
      Ala Arg Gly Tyr Asp Tyr Tyr Asp Ser Ser Gly Val Ala Ser Pro Phe
267
                                       105
269
      Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
```

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270
                                  120
              115
                                                       125
272 <210> SEQ ID NO: 21
273 <211> LENGTH: 333
274 <212> TYPE: DNA
275 <213> ORGANISM: Human
277 <400> SEQUENCE: 21
      cag tet gtg etg act cag eca ece tea geg tet ggg ace ece ggg cag
                                                                            48
      Gln Ser Val Leu Thr Gln Pro Pro Ser Ala Ser Gly Thr Pro Gly Gln
280
281
283
      agg gtc acc atc tct tgt tct gga agc acc tcc aac atc ggt act aat
                                                                            96
284
      Arg Val Thr Ile Ser Cys Ser Gly Ser Thr Ser Asn Ile Gly Thr Asn
285
                                        25
287
      act gca aac tgg ttc cag cag ctc cca gga acg qcc ccc aaa ctc ctc .
288
      Thr Ala Asn Trp Phe Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
289
291
      ate cae aat aat cag egg eee tea ggg gte eet gae ega tte tet
                                                                           192
292
      Ile His Asn Asn Asn Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser
293
                               55
295
      gge tee aag tet gge ace tea gee tee etg gee ate agt ggg ete cag
                                                                           240
296
      Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln
297
                                                75
299
      tet gag gat gag get gat tat tac tgt gea gea tgg gat gae age etg
                                                                           288
300
      Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ala Trp Asp Asp Ser Leu
301
303
      aat ggc cat tgg gtg ttc ggc gga ggg acc aag ctg acc gtc ctg
                                                                           333
304
      Asn Gly His Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
305
                  100
                                      105
307 <210> SEQ ID NO: 22
308 <211> LENGTH: 111
309 <212> TYPE: PRT
310 <213> ORGANISM: Human
312 <400> SEQUENCE: 22
314
      Gln Ser Val Leu Thr Gln Pro Pro Ser Ala Ser Gly Thr Pro Gly Gln
315
                                            10
317
      Arg Val Thr Ile Ser Cys Ser Gly Ser Thr Ser Asn Ile Gly Thr Asn
318
320
      Thr Ala Asn Trp Phe Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
321
323
      Ile His Asn Asn Asn Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser
324
                               55
326
     Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln
327
                           70
                                                75
329
      Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ala Trp Asp Asp Ser Leu
330
                                            90
332
      Asn Gly His Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
                                      105
335 <210> SEQ ID NO: 23
336 <211> LENGTH: 348
337 <212> TYPE: DNA
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VERIFICATION SUMMARY

DATE: 09/15/2004

PATENT APPLICATION: US/10/506,997

TIME: 16:21:21

Input Set : A:\SeqListing.txt

Output Set: N:\CRF4\09152004\J506997.raw

L:4 M:283 W: Missing Blank Line separator, <120> field identifier L:5 M:283 W: Missing Blank Line separator, <130> field identifier L:6 M:283 W: Missing Blank Line separator, <140> field identifier

L:6 M:270 C: Current Application Number differs, Replaced Current Application Number

L:7 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:12 M:283 W: Missing Blank Line separator, <160> field identifier